

DATE: 02/26/2003

1600

PATENT APPLICATION: US/09/804,733C TIME: 14:57:02 Input Set : D:\Mons016.app Output Set: N:\CRF4\02262003\I804733C.raw 3 <110 - APPLICANT: WANG, QI ET AL. 5 - 120 - TITLE OF INVENTION: RECOMBINANT PROTEINS CONTAINING REPEATING UNITS T 4:130 FILE FEFERENCE: MONS:016US 9 <140 - CUFRENT APPLICATION NUMBER: 09/804,733C C--> 10 <141> CURRENT FILING DATE: 2001-03-13 Does No Comply 12 -d150 - PRIOR APPLICATION NUMBER: PCT/US01/07957 Corrected Diskette Needec 13 -: 151 > PFIOR FILING DATE: 2001-03-13 15 - 160 - NUMBER OF SEQ ID NOS: 29 17 <170 - SOFTWARE: PatentIn Ver. 2.1 19 k210> SEQ ID NO: 1 20 K211 > LENGTH: 5 01 KO100 TYPE: PRT 22 - 213 - OFGANISM: Euthynnus pelamis 24 v400 / SEQUENCE: 1 15 Leu Lys Pro Ash Met 19 -0110- SEQ ID NO: 2 30 - 211 - LENGTH: 4 31 K212 - TYFE: FRT 32 - 0213 - OFGANISM: Euthynnus pelamis 34 -400 - SEQUENCE: 2 35 Lys Pro Ash Met 39 H210H SEQ IF NO: 3 40 HILLY LENGTH: 4 41 KILION TYPE: PRT 40 0213% OFGANISM: Euthynnus pelamis 44 ×400× SEQUENCE: 3 45 Val Val Tyr Ero 1 46 49 0210 SEQ ID NO: 4 50 k211 - DENGTH: 15 51 KILLS TYPE: DNA 52 - 211 - OFGANISM: Artificial Sequence 54 HODD - FEATURE: 55 - 2233 - OTHER INFOFMATION: Description of Artificial Sequence: Synthetic 0.6 Primer 58 HOLD FEATURE: 59 +333 * NAME/KEY: modified_base 60 <222 - LCCATION: (3)..(3) $61 \cdot 1223 \cdot \text{OTHER}$ INFORMATION: N = A, C, G or T/U 63 -: 4Ut - SEQUENCE: 4 15 W--> 64 ctnaarccna ayatg

RAW SEQUENCE LISTING

TATE: (2012)

TIME: 14:87:12

FATENT AFFLICATION: US/09/804,733C Input Set : D:\Mons016.app Cutput Set: N:\CRF4\02262003\1804733C.raw 67 K2105 SEQ ID NO: 5 68 <211 - LENGTH: +-0 69 K212 · TYFE: DNA 73 KB13 - CRGANISM: Artificial Sequence 71 H22 H FLATURE: 7: 322 OTHER INFORMATION: Description of Artificial Sequence: Synthetic 7.1 Frimer 7: :122 : FRATUPE: 70 - MARI - NAME/FEY: modified base 78 H22DH LOCATION: (3)..(54) 79 (20%) OTHER INFORMATION: N = A, C, G, or T/U81 K4000 SEQUENCE: 5 W--> 82 ctnaarccna ayatgctnaa rccnaayatg ctnaarccna ayatgctnaa rccnaayatg 60 85 H210+ SEQ IF NO: ϵ 85 H210+ LENGTH: 60 87 HILLS TYPE: DNA Se -00130 OFGANISM: Artificial Sequence 90 Fillor FEATURE: 91 KARAS OTHER INFOFMATION: Description of Artificial Sequence: Synthetic 91 Primer 94 -CACS FEATURE: 95 K2215 NAME/KEY: modified base 96 (1.1.1) LOCATION: (7). ((52)) (58) 97 $\pm 0.23 \pm 0.00$ OTHER INFORMATION: N = A, C, G or T/U 99 4400% SEQUENCE: 6 W--> 100 catrttnggy ttnagcatrt tnggyttnag catrttnggy ttnagcatrt tnggyttnag 60 103 K0105 SEQ ID NO: 7 164 K211 - LENGTH: 25 105 KUID TYPE: DNA 100 -0215 - ORGANISM: Artificial Sequence 10% -0.00 - FEATURE: 109 <2005 OTHER INFORMATION: Description of Artificial Sequence: Synthetic 110 Primer 110 KOIO - FEATURE: 113 MAME/KEY: modified_kase 114 <2220 LOCATION: (12)..(18) 115 <2233 OTHER INFORMATION: N = A, C, G or T/U117 <400 > SEQUENCE: 7 25 W--> 118 aaagaattcc tnaarccnaa yatgc 1.1 <01.05 SEQ ID NO: 8 100 +0010 LENGTH: 27 103 <010x TYPE: DNA 124 <213> ORGANISM: Artificial Sequence 126 KOBON FEATURE: 127 - COMES OTHER INFORMATION: Description of Artificial Sequence: Synthetic 12: Primer 130 KIZCH FEATURE: 131 HIRID MAME/KEY: modified base

RAW SEQUENCE LISTING

132 HODEL LOCATION: (18)..(24)



RAW SEQUENCE LISTING

FATENT AFFILICATION: US/09/804,733C

TATE: UP 16 Alle TIME: 14:87:02

27

Input Set : D:\Mons016.app

Output Set: N:\CRF4\02262003\I804733C.raw

133 \times 223 \times OTHER INFORMATION: N = A, C, G or T/U

135 4400 - SEQUENCE: 8

W--> 136 aaagcggccg ccatrttngg yttnagc

141 - 11 - TYPE: DNA

140 - . 13 - CR:ANISM: Artificial Sequence

144 - JOH - FEAFUFE:

145 - LLE - OTHER INFOFMATION: Description of Artificial Sequence: Synthetic

146 Frimer

144 4440 - SEQUENCE: 9

20 149 taatacgast cactataggg

15.1 - 210.4 SEQ IE NO: 10

153 - 211 - LENGTH: 19

154 ×21% - TYPE: DNA

155 - 113 - ORGANISM: Artificial Sequence

157 SILON FEATURE:

188 - 118 - OTHER INFOFMATION: Description of Artificial Sequence: Synthetic

Primer 159

161 < 400 > SEQUENCE: 10

19 160 ogateaatsa ogagrogee

165 - 2105 SEQ ID NO: 11

166 - 211 - LENGTH: 49

167 k212% TYPE: DNA

108 02135 ORGANISM: Artificial Sequence

170 - 100 - FEATUFE:

171 42.39 OTHER INFOFMATION: Description of Artificial Sequence: Synthetic

Frimer

174 ×120 + FEATUFE:

175 - III - MAME/KEY: rediffied_base

196 - 220 × LOCATION: (3)..(48)

177 - LLE - OTHER INFOFMATION: N = A, C, G or T/U

179 - 400 - SEQUENCE: 11

48 W--> 180 gtngtntayc cngtngtnta yccngtngtn tayccngtng tntayccn

183 k2105 SEQ ID NO: 12

194 - 211 - LENGTH: 48

185 KIIIP TYPE: DNA

186 - 213 - OEGANISM: Artificial Sequence

188 /2200 FEATURE:

189 <2235 OTHER INFORMATION: Description of Artificial Sequence: Synthetic

190 Primer

192 < 220 + FEATURE:

193 <2215 NAME/KEY: modified base

194 <222 > LOCATION: (1)..(46)

195 ± 213 OTHER INFORMATION: N = A, C, G or T/U

197 - 400 - SEQUENCE: 12

48 W--> 198 nggrtanach acnggrtana chachggrta nachachggr tanachac

201 - 110 - SEQ ID NO: 13

202 · F11 · LENGTH: 33

TATE: L. L. .

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TIME: 14:00: 2
                     PATENT APPLICATION: US/09/804,733C
                      Input Set : D:\Mons016.app
                      Output Set: N:\CRF4\02262003\I804733C.raw
     203 KLID - TYPE: DNA
     264 - 71: - ORGANISM: Artificial Sequence
     206 ·.L. · FEATURE:
     197 - . 1 - CTHER INFORMATION: Description of Artificial Sequence: Synthetic
              Frimer
     ::0 +:... FEATURE:
     ::1 +...! + NAME/WEY: modified_base
     1112 - ILL - LOCATION: (12)...33)
     113 - 113 - CTHEP INFORMATION: N = A, C, G or T/U
     LIB - 400 - SEQUENCE: 13
                                                                               33
W--> 216 aaaggatccg tngtntaycc ngtngtntay ccn
     119 - 110 - SEQ II NO: 14
     120 4.11 - LENGTH: 33
     DUI KUILA TYPE: INA
     [12] - 113 - OF GANISM: Artificial Sequence
     724 k270% FEATURE:
     115 WILLS OTHER INFOFMATION: Description of Artificial Sequence: Synthetic
              Frimer
     115 k200 - FEATUFE:
     P29 = 221 = NAME/REY: mcdified_base
     230 KISI - LOCATION: (10)..(31)
     231 + 23 + OTHER INFORMATION: N = A, C, G or T/U
     023 k400 > SEQUENCE: 14
                                                                                33
W--> 234 cccaagettn ggrtanaena enggrtanae nae
     037 ×210> SEQ ID NO: 15
     238 KA11 - LENGTH: 45
      134 - 11 - TYPE: DNA
     140 - 1130 OFGANISM: Artifleial Sequence
     242 A2209 FEATURE:
     143 - 123 - OTHER INFORMATION: Description of Artificial Sequence: Synthetic
     Frimer
     746 ×220> FEATURE:
     [47 . 2] AME (KEY: modified_base
     248 ×200 × LOCATION: (3)..(45)
     049 \times 103 \times \text{OTHER INFOFMATION: N = A, C, G or T/U}
      251 k4005 SEQUENCE: 15
W--> 252 gtnceneeng theeneengt neeneength ceneengthe encen
                                                                                45
      255 k210% SEQ ID NO: 16
      256 <211> LENGTH: 45
      157 HOLDH TYFE: DNA
      158 - 213 - OFGANISM: Artificial Sequence
     160 KILLOV FEATURE:
     061 4003> OTHER INFORMATION: Description of Artificial Sequence: Synthetic
     762
               Primer
      364 - 220 - FEATURE:
      165 + 121 > NAME/KEY: modified_base
      266 SELES LOCATION: (1)..(43)
      167 \cdot 113 \cdot \text{OTHER INFORMATION: N = A, C, G or T/U}
      269 -400> SEQUENCE: 16
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RAW SEQUENCE LISTING



TIME: 14:87:11 FATENT AFFLICATION: US/09/804,733C Input Set : D:\Mons016.app Cutput Set: N:\CRF4\02262003\I804733C.raw 45 W--> 270 nggnggnacn ggnggnacng gnggnacngg nggnacnggn ggnac 273 -010 - SEQ II NO: 17 274 - 11 - LENGTE: 36 275 - 31. - TYFE: DNA F . (RMANISM: Artificial Sequence 273 - J. D. FEATUFE: 279 - 1. : - CIHER INFORMATION: Description of Artificial Sequence: Synthetic Erim∈r 290 - . D. - FEATURE: LEB + DLT + NAME, FEY: medified base 284 -0220 - DOCATION: (12)...(36) LES - 113 - OTHER INFORMATION: N = A, C, G or T/U 280 - 400 - SEQUENCE: 17 36 W--> 288 aaaggateeg tneeneengt neeneengtn eeneen 191 - 110 - SEQ ID NO: 18 192 - 211 - LENGTH: 36 193 KININ TYPE: DNA 194 4213 + OFGANISM: Artificial Sequence 196 k210 + FEATURE: 197 (173) OTHER INFORMATION: Description of Artificial Sequence: Synthetic <u> 198</u> Frimer 300 - 320 - FEATURE: PG1 + LB1 + NAME/KEY: mcdified_base For which Location: (10)..(34) 303 + 203 + OTHEF INFOFMATION: N = A, C, G or T/U 305 - 400 - SEQUENCE: 18 36 W--> 306 aataagcttn ggnggnacng gnggnacngg nggnac 909 - 210 - SEQ ID NO: 19 310 - 211 - LENGTH: 8 311 - 212 - TYFE: PET 310 - 213 - OFGANISM: Artificial Sequence 314 - 200 - FEATURE: 315 ADDI: OTHER INFOFMATION: Description of Artificial Sequence:m Synthetic 37.47 Feptide 318 K4005 SEQUENCE: 13 319 Val Pro Pro Leu Lys Pro Asn Met 320 303 k010 SEQ ID NO: 10 304 k0115 LENGTH: 48 325 - DID - TYPE: DNA 3.6 x213 x OFGANISM: Artificial Sequence 328 + 220 + FEATURE: 329 <223 OTHER INFORMATION: Description of Artificial Sequence: Synthetic 330 Frimer

RAW SEQUENCE LISTING

930 AUDON FEATURE:

337 - 406 > SEQUENCE: 20

333 k221 + MAME/KEY: modified_base
334 k222 + MOCATION: (%)..(42)

135 < 0.35 OTHER INFORMATION: N = A, C, G or T/U



RAW SEQUENCE LISTING ERROR SUMMARY TATE: 14:50:00 FATERIT APPLICATION: US/09/804,733C TIME: 14:50:00

Input Set : D:\Mons016.app

Cutput Set: N:\CRF4\02262003\1804733C.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

```
Seg#:4; N Pos. 3,9
Seq#:5; N Pos. 3,9,18,24,33,39,48,54
Seq#:6; N Pos. 7,13,22,28,37,43,52,58
Seq#:"; N Pos. 12,18
Seq#:8; N Pcs. 18,24
Seq#:11; N Fcs. 3,6,12,15,18,04,27,30,36,39,42,48
Seq#:12; N Pos. 1,7,10,13,19,22,25,31,34,37,43,46
Seq#:13; N Pos. 12,15,21,24,27,33
Seq#:14; N Pas. 10,16,19,22,28,31
Seq#:15; N Pas. 3,6,9,12,15,18,11,24,27,30,33,36,39,42,45
Seq#:10; N Pos. 1,4,7,10,13,16,19,22,25,28,31,34,37,40,43
Seq#:17; N Pos. 12,715,18,21,24,17,30,33,36
Seq#:18; N Pcs. 10,13,16,13,22,25,28,31,34
Seq#:20; N Pcs. 3,6,9,12,18,27,20,33,36,42
Seq#:21; N Pcs. 7,13,16,14,22,31,37,40,43,46
Seq#:22; N Pcs. 13,16,14,22,31,37,40,43,46,52
Seq#:23; N Pcs. 19,25,28,31,34,40,43,46,85,61,64,67,70,76,79,82
Seq#:05; M Pcs. 3,6,9,15,18,.1,14,30,39,41,45,51,54,57,60,66
Seq#:26; N Pos. 7,13,16,19,22,28,31,34,43,49,52,55,58,64,67,70
Seq#:27; N Pos. 13,16,19,85,23,31,34,40,49,52,55,61,64,67,70,76
Seq#:28; N Pos. 19,25,28,31,34,40,43,46,55,61,64,67,70,76,79,82
```